

OTPE



RAW SEQUENCE LISTING

US/10/077,176 PATENT APPLICATION

DATE: 03/04/2002 TIME: 14 46:32

Input Set : A:\433480_1.txt

Output Set: N:\CRF3\03042002\J077176.raw

- 4 <110 APPLICANT: Brachmann, Painer
- 6 <120° TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR 253
- 9 <130 FILE PEFERENCE: 004255.00008
- C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/077,176
- C--> 11 <141> CURRENT FILING DATE: 2002-02-19
 - 11 <160 · NUMBER OF SEQ ID NOS: 71
 - 13 <170 SOFTWARE: FastSEQ for Windows Version 4.0
 - 15 -: 210 SEQ ID NO: 1
 - 16 -211 LENGTH: 1182
 - 17 212 TYPE: DNA
 - 18 <213 · ORGANISM: Artificial Sequence
 - 20 RU20 FEATURE:
 - 21 < 223 % OTHER INFORMATION: Produced by genetic engineering
 - 23 < 490 + SEQUENCE: 1
 - 24 atyguagaan cacagtoaga tootagegto gaaccaccto tgagtoagga aaccttttoa 60 25 gaddigtgga aattgcttoo tgaaaacaac gttotgtooc cattgcctag tcaagcaatg 120 180
 - 26 gatgatttga tyctytocco agacgatatt gaacaatggt toactgaaga tocaggooca 27 gatgaagete caegaatgee agaggeeget ecaeeggttg eeceageaee ageagetest 240 300
 - 28 acadeggegg deceagates ggoodcated tggootetgt catottotgt continuous
 - 29 aaaaootaco agggoagota eggttteegt etgggettet tgeattetgg aaetgecaag
 - 30 totgttactt glacgiacte tecagecett aacaagaigt titgecaact egegaagaee
 - 31 typocagted aactgtgggt cgactccacc cotocacctg gtacacgtgt cogcgcaatg
 - 32 gecatetaca ageagageea geacatgaeg gaggtegtae gaegetgtee acaecatgag
 - 33 cgctgctcag attctgatgg tctggcgcca ccacagcatc ttatccgagt ggaaggtaac 34 ctacgcgtgg agtatetaga tgacegeaac aettttegae acagtgtggt ggtgeeatat
 - 35 gaggoaccag aagttygote tgactgoacc accatcoact acaactatat gtgtaacagt
 - 36 tcatgcatgg gcggcatgaa ccggcggccg atcctgacca tcatcactct cgaggattcc
 - 37 trangtaate tectaggang gaatteettt gaggtgegtg titgtgeatg oneggggenge
 - 38 gategeegga eegaagagga gaateteegg aagaaaggtg ageeteacea egagetgeea
 - 39 ccauquagea ctaagegage actgeeaaac aacaccagea gttetecaea gecaaagaag
 - 40 aaasetttgg aeggagaata titeaeeett eagateegtg geegtgageg gitegagatg
 - 41 ttorgagage tgaatgagge ettagaaett aaggatgeee aggetggtaa ggageeagga
 - 42 ggcagoogtg ctcatagoag coacotgaag tocaaaaagg gtcagtctac etecogocat
 - 43 aaaaaactga tgttcaagac cgaaggtcot gactcagact ga
 - 45 -2100 SEQ ID NO: 2
 - 46 211: LENGIH: 1182
 - 47 PILLS TYPE DNA
 - 48 213 ORGANISM. Artificial Sequence
 - 50 <21): FEATURE:
 - OTHER INFORMATION: Produced by genetic engineering 51×223
 - 53 <400> SEQUENCE: 2
 - 54 atgjaagaac cacagtcaga tootagogto qaaccaccoo tqagtcagda aaccttttca
 - 55 gatotytyga agottottoo tgaagacaac gttotytooc cattyootag teaagcaatg

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RAW SEQUENCE LISTING IAIE: 93/34,2002
PATENI APPLICATION: US/10/077,176 IIME: 14 46 32

Input Set : A:\433480_1.txt

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56 datigatitua tyotigagooo adacqatat, qaabaatigut toactdadga tucadgood	leč
carrattor adaddcodct coacciditu tercadeass adeada	24
appointed discrete fideblication and colored in the colo	130
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- 11 1 antique point at another and colour decorated and and an antique of the colour section and the colour s	480
and the second second description and the second se	540
- va	600
- v. lilining variational transcription delicities delicited according to Maratina and	660
and the second constitution of the original according a conduction of the second secon	720
- January Control to a control of the state	734
	840
- /b-saaraa saaraaraa daafofdodi Badadadquuq aybbbbasaa saaraa saaraa	900
and the same of the same and the same of t	960
7) aaacctttgg acggagaata tttcaccctt cagatccgtg geogtgageg gttcgagatg	1020
71 ttocqaqago tgaatgaggo ottagaactt aaggatgooc aggotggtaa ggagecagga	1080
7) trecquadge tyaatyagge betagaabb aby trecquadate greater creecqueat 72 ageagoogtg ofcatageag coacetgaag tecaaaaaagg greagtetac creecqueat	1140
73 aaaaaastga tyttcaagac cgaaggtoot gactcagact ga	1132
7) aaaaaactga tyttcaayac cyuuyyesee yuuushin j	
75 - 21(c) SEQ ID NO: 3	
76 - 211: LENGTH: 1181	
77 - 2110 TYPE: DNA	
78 - 213 - OFGANISM: Artificial Sequence	
80 (120) FEATURE: 81 (123) OTHER INFORMATION: Produced by genetic engineering	
83 (400) SEQUENCE: 3 84 atggaagaac cacagtoaga tootagogto gaaccacoco tgagtoagga aaccttttoa	$\epsilon \circ$
of the state of the translation of the state	120
anacqatatt qaaqqqq xqqqq aqacqatatt qaaqqqq cca, cqaqqq	180
and the many of the same and th	24(.
88 aladeagong coccagetoe ageoconates transfer transfering aactgoonag	3(:()
88 araceggedg coebagette ggottlede oggettet tgeattetgg aactgedaag 89 aaaacetace agggeageta eggtteegt etgggettet tgeattetgg aactgedaag	360
The second second transfer of the contract of	420
91 typopagtop aactgtgggt cgactopace octopacetg gtapacqtgt regognaatg	480
92 gocatitaca agcagagoca goacatgacg gaggtoqtac gacqctqtoc acaccatgag	540
93 controlled attempting totgogode coacaycate that coagt yeargetaac	600
93 edetigetedg alleetgatgg setgyngdda seattitegae acagtgtggt ggtgeeatat 94 etaegegtgg agtatetaga tgaeegeaae actititegae acagtgtggt ggtgeeatat	660
94 ctacgograf aglatolaga tyaccycado accatecact acaactatat gtgtaacagt 95 gagccaccag aagttggcto tgactgcacc accatecact acaactatat gtgtaacagt	720
96 teatquatgg geggeatgaa deggeggeeg atectgalea teateactet egaggattee	780
66 thatquatg geggeargan cegggggggggggggggggggggggggggggaggaggagga	840
67 thagginate tectaggacy gaateboote gagsas gits ageotherea egagetyeea 68 gategoogya cogaagagga gaatetoogg aagaaa ggtg ageotocca egagetyeea	900
63 gatogoogga cogaagagga gaatoooogg dagaacgagot ottotooaca gocaaagaag	960
100 aaaaatttag acgagaata ttteacectg cagattegtg geogtgageg gttegagatg	1020
100 aaacetttag acggagaata tittaacetta cagtatacea aggetgataa gaagecagga 101 ticcuagage tgaatgagge ettagaacet aaggatgeca aggetgataa tagtatacat	1030
101 thochagago tigaatgaggo obtagaadte daggadagada garagoonga 101 qqcaqooggg occattogto toacotgaag tocaaaaaagg gtoagtotac tagtogocat	1110
101 ggcagergag cocattegue teactigady esteadadagy goody	1181
102 aaaaaactga gttcaagacc gaaggtcotg actcagactg a	
105 42100 SEQ ID NO: 4	
16(<211> LENGTH: 1182	
16" <212> TYPE: INA	
108 <213> ORGANISM: Artificial Sequence	

RAW SEQUENCE LISTING

TAIE: 3/04/2002 FATENI APPLICATION: US/10/077,176 IME: 14:44:32

Input Set : A:\433480_1.txt

output Set: N:\CRF3\03042002\J077176.raw

110 ×220 FEATURE:	
111 - 223: OTHER INFORMALLON: Produced by denetic engineering	
A CONTRACT OF THE PROPERTY OF	60
113 k40 : SEQUENCE: 4 114 atquaaqaac cacagtcaga teetageqte qaancacete tgagtegaga aacettitea	120
The state of the same of the s	180
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The state of the s	420
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	7 20
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and the second second destruction and design and described and the second secon	960
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- and the common transfer of the manager and additional	1140
13) greatered of catagoay coacctgaag tocaddadyy greatered chocky the	1182
133 maaaaactga tgttcaagac cgaaggtoot gactcagact ga	TICE
135 - 2105 SEQ ID NO: 5	
136 + 211> LENGTH: 1181	
137 (212) TYPE: UNA	
138 -2213 OFGANISM: Artificial Sequence	
1 A TO TO GO OF TEATING .	
14) - 270. Thatoke: 141 - 223: OTHER INFORMATION. Produced by genetic engineering	
A CONTRACT F	60
and the same a contrargat contaggator aaccacctct gagtcaggad acculturage	120
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- I I I I I I I I I I I I I I I I I I I	1050
162 gragoratge tratagrade caretyaagt craaaaaggg tradtetace terrepresata	1140
102 4044004040 404042	

RAW SEQUENCE LISTING

TATE: 017.4721.4 PAIENT APPLICATION: US/10/077,176 IIME: 14:40:32

Input Set : A:\433480_1.txt
Output Set: N:\CRF3\03042002\J077176.raw

	1161
163 augaactyat yttoaayaee yaayyteetu aeteayaetu a	LICI
165 k210> SEQ ID NO: 6	
leb <211> LENGTH: 1182	
167 K212 TYPE: DNA	
168 <113 + ORGANISM: Artificial Sequence	
37 (2) OO. FEATURE:	
171 8223 OTHER INFORMATION Produced by genetic engineering	
A TO A CENTENCE 6	
and the same and anatomical techniques and adolescent the same and the	60
- the contract the translation of the contract	120
- I I I I I I I I I I I I I I I I I I I	180
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and the second of the second o	430
TO A CONTRACT MANUAL MA	540
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and the second s	780
and the tracks tracks are a description of the desc	840
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and a contract of a graduation and a carried	960
the carrier of the carrier and the carrier of the c	1020
and the second transfer and office of the second additional distriction of the second	1030
100 aggarageta etcatageau ceacetgaag tecadadagg geoagtetae etcatageau	114
192 agaawactga tgttcaagas cgaaggtcot gactcagact ga	1182
195 - 210 - SEQ ID NC: 7	
196 - 211 - LFNGTH: 1182	
197 - 212 - TYPE: DNA	
198 - 113 - OFGANISM: Artificial Sequence	
CONTRACTOR DISTRIBUTE	
200 - 220 - FRATORM. 201 - 223 - OTHER INFORMATION: Produced by genetic engineering	
A SA A GREAT CONTRACT A CONTRACT	
and the same appropriate and the tage of the dade of the tage of tage	60
and the state of t	120
a contract of the contract of	180
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216 traigeating graduated being consisted to training and additional strangers and all training and additional straightful and the straightful and	840
21 Codyquated Cooperage of Section 18 As a section of the section	

TATE: 03/04/2002 TIME: 14:46:32 RAW SEQUENCE LISTING PATENT APPLICATION: US/10/077,176

Input Set : A:\433480_1.txt

matput Set: N:\CRF3\03042002\J077176.raw

autput Set: N:\CRF3\03042002\J07/1/6.raw	
216 dategoogga eegaayagga gaateteegg aaraaaadutg arestereega edagotgooga 216 dategoogga eegaayaaa	<u>(4)</u>
nataronia comandada gaatotocar aaraaadati datta roaga recaaadaad	960
lle dategeogga cegaagaga gaatemeeri aadadaga utteteeaga decaaagaagaagaagaagaagaagaagaagaagaagaagaa	1020
21) coaddaadea staaqeqada actgocaaac aaraccadea geterbada uttegagata 21) coaddaacttag acggagaata titeaccett cadatecdid deegidaacg uttegagata 22) aaarettiag acggagaata titeaccett aaddatgeee adgetgataa ggaggeadda	1080
221 tiredagage tgaatgagie ettagaaett aaggatgee aggetgetaa ggageeaat 221 tiredagage tgaatgagie eracetgaag tecaaaaagg dicadtetae eteoogocat	1140
221 ticedagage tyantyagaa ccaectyaay tecaaaaagg dicadistae ecologodae	1182
221 tinedagage tgaatgagge ettadaaett aaddatgeed addieggeda etebogoest 221 geadeegtg eteatageag eeacetgaag teeaaaaagg dieadtotae etebogoest 222 ggaadeegtg eteatageag egaaggieet gaeteagaet ga 223 gaaggaantga tgitteagae egaaggieet gaeteagaet ga	
223 walaaantaa tottoaayaa caasaya	
225 (210; SEQ ID NG: 8	
226 <2115 FNGTH: 1182	
22 * (2) LYPE DNA	
22' (212) HYPE DAS 228 (213) (FGANISM) Artificial Sequence	
230 <220 FEATURE 231 <223 - THEE INFORMATION: Produced by genetic engineering	
231 < 223 · OTHER INFORMATION: PROGRESS 4 Dy	60
223 & 100 SEQUENCE: 8	-
233 (4)0% SEQUENCE: 8 234 atgguaguad dadagtdaga tedtagegto daaddastte tgagtdagga adscrittica 234 atgguaguad dadagtdaga tedtagegto daaddastte cattgedtag tedaggastag 235 qaddtgtgga aattgettee tgagaaadaad gttotgtood cattgedtag tedaggooda	120
nas gardinida aditiquide egalari	130
ose datastitude indirection against a second agreement against additioned	24:)
non material cacquaryce agains	3()()
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216 Estatiacti didoquades essenti confidences	480
211 tarceagtee adolytyy	540
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out date and the additional control of the control	780
51. +31.4:4:4:4:4:4:4:4:4:4:4:4:4:4:4:4:4:4:4:	840
ner and the following yours and a second to the control of the con	9 (14)
24 gatogoogga gogaaqagga gaatotooyg aagaaaggtg agocataagaagaag	960
24 thingshig a cogaagaga gaateteeyg aagaanggig ageetadaa geeaaagaaga 249 gategoogga cogaagagga actgooaaac aacaccagca gttotucaca geeaaagaaga 249 ccaqgaagca ctaagogagc actgooaaac aacaccagca gttotucaca geeaaagaaga	1020
249 chaqqaaqca ctaaqcqaqc actqccaaac aacaccaqca gutuulaada qutucqaqatq 250 aaacctttgg acggaqaata tttcaccctt caqatccqtq googtqaqcq qttcqaqaq 250 aaacctttgg acggaqaata tttcaccctt aaqqatgccb aggctqqtaa gqaqccaqqa	1080
250 alacettigg acggagaata titteaceett cagaticegti geografiga gyagecagga 250 titeegajage tigaatigagge eitagaaett aangatgeet aggetigtaa gyagecagga 250 titeegajage tigaatigagge esacetigaag teeaaaaagg gicagtetae eiceegeeat	1140
25% data da	1182
252 ggcageogtg etcatagody classesyddy gaetcagaet ga 253 gaaagaetga tgtteaagae egaaggteet gaetcagaet ga	
153 dadadactyd tyddau 9 153 - 210: SEQ ID NO: 9	
256 - 211 LENGTH: 1182	
total tot	
>	
2:8 21: OFGANISM: ARCTITICIAL Sequence of the	
260 - 210 - FEATUPE: 261 - 223 OTHER INFORMATION. Produced by genetic engineering	
2.1 223 WIRE THE CONTROL OF THE CONT	€ 0
263 (400) SEQUENCE: 9 264 159944 again cacagtoaga tootagouto gaaccacoto tqaqtoagga adoptittoa 264 159944 gaac cacagtoaga tootagouto gaaccacoto tqaqtoagga adoptittoa	120
263 Programmes cacaghoaga toctagosto gaaccacolo tyaqromaya 264 Programmas aattgottoo tyaaaacaac gitotgtooc cathgochag ichaggaaatg 265 qalotsitgaa aattgottoo tyaaaacaac gitotgtooc cathgochag tolaaggocca	180
264 1943444424 aattgettee tgaaaacaac gitetgieed eatigeties 265 qaictittga aattgettee agacgatatt gaacaatggt teactgaiga teraggeeet 266 qaigaitttga tgetgieed agacgacget coaceggitg coccagoide agacgacteet	240
266 gatgatttga tgctgtcccc agacgatatt gaacaatggt toachgarga agcagotect 267 datgatgtgctc bacgaatgcc agaggccqt coaceggttg coccupate agcagotect 267 datgatgctc bacgaatgcc agaggccqtcc tigoctctgt catottctgt coettcccag	300
2.7 dangligete bacquatqce agaggeodet chaecqquiq coolings bettercag 2.7 dangligete bacquatqce qqccccatec tigoctetqt catonicigt chettercag 2.63 abaccqqcqq ecceagete qqccccatec tigoctetqt tigoattetqq dactqbcaaq	360
263 acidotace aggreageta egytttengt otgacetat tipostotig aantginaag 269 aanandtace aggreageta egytttengt otgacetat titostotig aantginaage 269 aanandtace aggreageta tonagheett andaagatat titagenaact eninganaatg	420
269 aalacetace agggeageta eggttteegt etgadetist sychiseact elegalgace 270 tetetactt graegiacte telagocate aleaagatgt titigeraact elegalgact segentaatg	480
270 totottactt gracquacto totagocott alcaagatgt betgraden totagogoaatg 271 tooccagtoc aactgtgggt cqactocacc cotocacctg gracatgtoc acaccatgag	540
271 todocagtoc aactgtgggt cqactdoacd cotdoacdty gtucusary 271 todocagtoc aactgtgggt cqacatgacg gaggtcytac gacgstgtoc acaccatgag 272 qocatctaca agcagagcca qcacatgacg gaggtcytac gacgstgtoc acaccatgag	w
272 godatotada ayuayayaan arr	and in the
nas peen	islected in the

gse of this is on waa has been detected in the Sequence Listing Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence ising n or X.aa

VERIFICATION SUMMARY

1A1E: 13/04/2004 IIME: 14:46:33 PATENT APPLITATION: US/10/077,176

Input Set : A:\433480_1.txt

entrut Set: N:\CRF3\03042002\J077176.raw

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Lill MilTo C: Current Application Number Hitlers. Replaced Jurrent Application N
1:11 M:271 C: Current Filing Date differs. Replaced Current Filing Date
L:2328 M:341 W: (46: "n" or "Xaa" used, for SEQ ID#:71
1:2332 M 341 W: (46: "n" cr "Xaa" used. for SEQ ID=:71
L:2336 M.341 W: (46: "n" cr "Xaa" used. for SEQ ID=:71
L:2340 M:341 W: (46: "n" cr "Xaa" used. for SEQ ID=:71
L 2344 M.341 W: (46) "n" or "Xaa" used, for SEQ ID=-71
L:2343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID= 71
1:2352 M:341 W: (4t) "n" cr "Xaa" used, for SEQ ID#-71
L:2355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#.71
L:2360 M:341 W: (46) "n" (1 "Xaa" used, for SEQ ID=:71
L:2364 M:341 W: (4r) "n" or "Xaa" used, for SEQ ID::71
L 2368 M:341 W: (4+) "n" or "Xaa" used, for SEC ID::71
L:2372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID=:71
L:2376 M:341 W: (40) "n" or "Xaa" used, for SEQ ID::71
L:2380 M:341 W: (4+) "n" or "Xaa" used, for SEQ ID#:71
L:2384 M:341 W: (4b) "n" or "X:a" used, for SEQ ID#:71
L:2388 M:341 W: (4b) "n" or "Xaa" used, for SEQ ID#:71
 L:1392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
 L:1396 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
 L:24(00 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
 L:2404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
 L:2408 M:341 W: (46) "n" or "Xaa" used, for SEg ID#:71
 L:2408 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
 L:2420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
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